'01/0 46	00" adj base adj EPO; JPO; DERWENT	"14000" pairs	L14 13	BRS 1	12
PGPUB; 2003/01/0 2 17:46	USPAT; US- 12 EPO; JPO; DERWENT	9 same	L13 0	BRS 1	11
PGPUB; 2003/01/0 2 17:45	0" adj base adj EPO; JPO; DERWENT	"13000" pairs	L12 5	BRS 1	10
PGPUB; 2003/01/0 2 17:41	USPAT; US- 10 EPO; JPO; DERWENT	l 9 same	L11 71	BRS 1	٥
PGPUB; 2003/01/0 2 17:40	uspat; us- adj pairs EPO; JPO; DERWENT	42328 base	L10 42	BRS 1	ω
PGPUB; 2003/01/0 2 17:39	USPAT; US- 5 same 7 EPO; JPO; DERWENT	95 4 same	L9 65	BRS 1	7
PGPUB; 2003/01/0 2 17:45	3" adj base adj EPO; JPO; DERWENT	"12793" pairs	L8 1	BRS 1	O
PGPUB; 2003/01/0 2 17:38	brate or EPO; JPO; DERWENT	48103 (verte	L7 48	BRS 1	Л
PGPUB; 2003/01/0 2 17:37	USPAT; US-F EPO; JPO; DERWENT	11445 exon	L5 11	BRS]	4
PGPUB; 2003/01/ 2 17:36	uspar; us- adj acid EPO; JPO; DERWENT	95984 nucleic	L4 95	BRS]	ω
PGPUB; 2003/01/0 2 17:36	USPAT; US- EPO; JPO; DERWENT	spastin	L3 6	BRS]	Ν
PGPUB; 2003/01/0 2 17:28	USPAT; US- EPO; JPO; DERWENT	5 arsacs	L1 85	BRS]	Ъ
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FILE 'HOME' ENTERED AT 17:51:15 O
=> file medline caplus biosis embase scisearch agricola
                                                 SINCE FILE
                                                                  TOTAL
COST IN U.S. DOLLARS
                                                       ENTRY
                                                                SESSION
FULL ESTIMATED COST
                                                        0.21
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=> s arsacs
            56 ARSACS
L1
=> s spastin
           174 SPASTIN
=> s nucleic acid (p) exon (p) (vertebrate or human)
   3 FILES SEARCHED..
           724 NUCLEIC ACID (P) EXON (P) (VERTEBRATE OR HUMAN)
=> s 13 (p) 12
             0 L3 (P) L2
=> s 12 (p) (human or vertebrate) (p) gene
   3 FILES SEARCHED...
            43 L2 (P) (HUMAN OR VERTEBRATE) (P) GENE
=> duplicate remove 15
DUPLICATE PREFERENCE IS 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
PROCESSING COMPLETED FOR L5
             33 DUPLICATE REMOVE L5 (10 DUPLICATES REMOVED)
=> s 16 (p) (base pair)
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L43 (P)
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L45 (P) '
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L47 (P) '
             0 L6 (P) (BASE PAIR)
L7
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     (FILE 'HOME' ENTERED AT 17:51:15 ON 02 JAN 2003)
     FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT
     17:51:42 ON 02 JAN 2003
             56 S ARSACS
            174 S SPASTIN
L2
L_3
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=> s 16 (p) exon
PROXIMITY OPERATOR LEVEL NOT CONSIDERT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L56 (P) EXON'
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L58 (P) EXON'
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L60 (P) EXON'
             1 L6 (P) EXON
=> d 18 1 ibib abs
     ANSWER 1 OF 1 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
                    2000:350945 BIOSIS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                    PREV200000350945
                    Clinical and pathologic findings in hereditary spastic
TITLE:
                    paraparesis with spastin mutation.
                    White, K. D.; Ince, P. G.; Lusher, M.; Lindsey, J.;
AUTHOR (S):
                    Cookson, M.; Bashir, R.; Shaw, P. J.; Bushby, K. M. D. (1)
CORPORATE SOURCE:
                    (1) Department of Human Genetics, 19/20 Claremont Place,
                    Newcastle upon Tyne, NE2 4AA UK
                    Neurology, (July 12, 2000) Vol. 55, No. 1, pp. 89-94.
SOURCE:
                    print.
                    ISSN: 0028-3878.
DOCUMENT TYPE:
                    Article
                    English
LANGUAGE:
SUMMARY LANGUAGE:
                    English
     Objective: To describe a family with chromosome 2p-linked hereditary
     spastic paraparesis (HSP) associated with dementia and illustrate the
     cerebral pathology associated with this disorder. Background: HSP
     comprises a heterogeneous group of inherited disorders in which the main
     clinical feature is severe, progressive lower limb spasticity. Nongenetic
     classification relies on characteristics such as mode of inheritance, age
     at onset, and the presence or absence of additional neurologic features.
     Several loci have been identified for autosomal dominant pure HSP. The
     most common form, which links to chromosome 2p (SPG4), has recently been
     shown to be due to mutations in spastin, the gene encoding a novel
     AAA-containing protein. Results: The authors report four generations of a
     British family with autosomal dominant HSP in whom haplotype analysis
     indicates linkage to chromosome 2p. In addition, a missense mutation has
     been identified in exon 10 of the spastin gene (A1395G). Dementia was
     documented clinically in one member of the family, two other affected
     family members were reported to have had late onset memory loss, and a
     younger affected individual showed evidence of memory disturbance and
     learning difficulties. Autopsy of the demented patient confirmed changes
     in the spinal cord typical of HSP and also demonstrated specific cortical
     pathology. There was neuronal depletion and tau-immunoreactive
     neurofibrillary tangles in the hippocampus and tau-immunoreactive balloon
     cells were seen in the limbic and neocortex. The substantia nigra showed
     Lewy body formation. The pathologic findings are not typical of known
     tauopathies. Conclusions: The authors confirm that chromosome 2p-linked
     HSP can be associated with dementia and that this phenotype may be
     associated with a specific and unusual cortical pathology.
=> d his
     (FILE 'HOME' ENTERED AT 17:51:15 ON 02 JAN 2003)
     FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT
     17:51:42 ON 02 JAN 2003
             56 S ARSACS
L1
            174 S SPASTIN
L2
            724 S NUCLEIC ACID (P) EXON (P) (VERTEBRATE OR HUMAN)
L3
L4
              0 S L3 (P) L2
L5
             43 S L2 (P) (HUMAN OR VERTEBRATE) (P) GENE
L6
             33 DUPLICATE REMOVE L5 (10 DUPLICATES REMOVED)
              0 S L6 (P) (BASE PAIR)
L7
              1 S L6 (P) EXON
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